

05081147CRF
SEQUENCE LISTING

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FUCHS, Robert, Pierre, Paul
OLIERIC, Vincent

<120> PROTEIN CRYSTAL COMPRISING THE PROCESSIVITY CLAMP FACTOR
OF DNA POLYMERASE AND A LIGAND, AND ITS USES

<130> 0508-1147

<140> US 10/561,867

<141> 2006-07-06

<150> PCT/EP2004/006942

<151> 2004-06-25

<150> EP 03291596.9

<151> 2003-06-27

<160> 7

<170> PatentIn version 3.5

<210> 1

<211> 16

<212> PRT

<213> Escherichia coli

<400> 1

Val Thr Leu Leu Asp Pro Gln Met Glu Arg Gln Leu Val Leu Gly Leu
1 5 10 15

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<212> PRT

<213> Artificial sequence

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<223> synthetic peptide

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Arg Pro Val Lys Val Thr Pro Asn Gly Ala Glu Asp Glu Ser Ala Glu
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Ala Phe Pro Leu Glu Phe
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<210> 3

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<212> DNA

<213> Artificial sequence

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<223> synthetic oligonucleotide - Primer for replication assay
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gtaaaacgac ggccagtgcc aagcttagtc 30

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<211> 90
<212> DNA
<213> Artificial sequence

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<223> synthetic oligonucleotide - template for replication assay

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ccatgattac gaattcagtc atcaccggcg ccacagacta agcttggcac tggccgtcgt 60
tttacaacgt cgtgactggg aaaaccctgg 90

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<212> PRT
<213> Escherichia coli

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Met Lys Phe Thr Val Glu Arg Glu His Leu Leu Lys Pro Leu Gln Gln
1 5 10 15

Val Ser Gly Pro Leu Gly Gly Arg Pro Thr Leu Pro Ile Leu Gly Asn
20 25 30

Leu Leu Leu Gln Val Ala Asp Gly Thr Leu Ser Leu Thr Gly Thr Asp
35 40 45

Leu Glu Met Glu Met Val Ala Arg Val Ala Leu Val Gln Pro His Glu
50 55 60

Pro Gly Ala Thr Thr Val Pro Ala Arg Lys Phe Phe Asp Ile Cys Arg
65 70 75 80

Gly Leu Pro Glu Gly Ala Glu Ile Ala Val Gln Leu Glu Gly Glu Arg
85 90 95

Met Leu Val Arg Ser Gly Arg Ser Arg Phe Ser Leu Ser Thr Leu Pro
100 105 110

Ala Ala Asp Phe Pro Asn Leu Asp Asp Trp Gln Ser Glu Val Glu Phe
115 120 125

Thr Leu Pro Gln Ala Thr Met Lys Arg Leu Ile Glu Ala Thr Gln Phe
130 135 140

Ser Met Ala His Gln Asp Val Arg Tyr Tyr Leu Asn Gly Met Leu Phe
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155

145		150		160
Glu Thr Glu Gly	Glu Glu Leu Arg Thr Val	Ala Thr Asp Gly	His Arg	
	165	170	175	
Leu Ala Val	Cys Ser Met Pro Ile	Gly Gln Ser Leu Pro	Ser His Ser	
	180	185	190	
Val Ile	Val Pro Arg Lys Gly	Val Ile Glu Leu Met	Arg Met Leu Asp	
	195	200	205	
Gly Gly	Asp Asn Pro Leu Arg	Val Gln Ile Gly	Ser Asn Asn Ile Arg	
	210	215	220	
Ala His	Val Gly Asp Phe Ile Phe Thr	Ser Lys Leu Val	Asp Gly Arg	
	225	230	235	240
Phe Pro	Asp Tyr Arg Arg Val	Leu Pro Lys Asn Pro	Asp Lys His Leu	
	245	250	255	
Glu Ala	Gly Cys Asp Leu Leu Lys	Gln Ala Phe Ala Arg	Ala Ala Ile	
	260	265	270	
Leu Ser	Asn Glu Lys Phe Arg	Gly Val Arg Leu Tyr	Val Ser Glu Asn	
	275	280	285	
Gln Leu	Lys Ile Thr Ala Asn Asn Pro	Glu Gln Glu Glu Ala Glu Glu		
	290	295	300	
Ile Leu	Asp Val Thr Tyr Ser Gly Ala	Glu Met Glu Ile Gly Phe Asn		
	305	310	315	320
Val Ser	Tyr Val Leu Asp Val Leu Asn	Ala Leu Lys Cys Glu Asn Val		
	325	330	335	
Arg Met	Met Leu Thr Asp Ser Val	Ser Ser Val Gln Ile Glu Asp Ala		
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Ala Ser	Gln Ser Ala Ala Tyr Val	Val Met Pro Met Arg Leu		
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 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthetic peptide - derived from SEQ ID NO 1

<400> 6

Arg Gln Leu Val Leu Gly Leu
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<210> 7

<211> 210

<212> PRT

<213> Artificial Sequence

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<223> synthetic peptide - derived from beta clamp of
E. coli DNA polymerase III

<400> 7

Leu Asn Gly Met Leu Phe Glu Thr Glu Gly Glu Glu Leu Arg Thr Val
1 5 10 15Ala Thr Asp Gly His Arg Leu Ala Val Cys Ser Met Pro Ile Gly Gln
20 25 30Ser Leu Pro Ser His Ser Val Ile Val Pro Arg Lys Gly Val Ile Glu
35 40 45Leu Met Arg Met Leu Asp Gly Gly Asp Asn Pro Leu Arg Val Gln Ile
50 55 60Gly Ser Asn Asn Ile Arg Ala His Val Gly Asp Phe Ile Phe Thr Ser
65 70 75 80Lys Leu Val Asp Gly Arg Phe Pro Asp Tyr Arg Arg Val Leu Pro Lys
85 90 95Asn Pro Asp Lys His Leu Glu Ala Gly Cys Asp Leu Leu Lys Gln Ala
100 105 110Phe Ala Arg Ala Ala Ile Leu Ser Asn Glu Lys Phe Arg Gly Val Arg
115 120 125Leu Tyr Val Ser Glu Asn Gln Leu Lys Ile Thr Ala Asn Asn Pro Glu
130 135 140Gln Glu Glu Ala Glu Glu Ile Leu Asp Val Thr Tyr Ser Gly Ala Glu
145 150 155 160Met Glu Ile Gly Phe Asn Val Ser Tyr Val Leu Asp Val Leu Asn Ala
165 170 175Leu Lys Cys Glu Asn Val Arg Met Met Leu Thr Asp Ser Val Ser Ser
180 185 190

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Val	Gln	Ile	Glu	Asp	Ala	Ala	Ser	Gln	Ser	Ala	Ala	Tyr	Val	Val	Met
		195					200					205			

Pro	Met
	210